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BIOTECHNOLOGY
SYSTEMS
BRANCH

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/67/,773
Source: 01PE
Date Processed by STIC: 6/11/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:
  - U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
  - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 01/29/2002

JUL 1 2002



OIPE

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RAW SEQUENCE LISTING
                                                             DATE: 06/11/2002
                     PATENT APPLICATION: US/09/671,773
                                                             TIME: 14:12:20
                     Input Set : A:\PTO.VSK.txt
                     Output Set: N:\CRF3\06112002\1671773.raw
                                                                      Does Not Comply
                                                                  Corrected Diskette Needed
      3 <110> APPLICANT: BAYER CORPORATION
      5 <120> TITLE OF INVENTION: Pituitary Adenylate Cyclase Activating Peptide (PACAP)
 Receptor 3
              (R3) Agonists and Their Pharmacological Methods of Use
      .6
      8 <130> FILE REFERENCE: Bayer
     10 <140> CURRENT APPLICATION NUMBER: US/09/671,773
   > 10 <141> CURRENT FILING DATE: 2002-05-31
     10 <150> PRIOR APPLICATION NUMBER: 09/407,832
     11 <151> PRIOR FILING DATE: 1999-09-28
     13 <150> PRIOR APPLICATION NUMBER: 09/595,280
     14 <151> PRIOR FILING DATE: 2000-06-15
     16 <160> NUMBER OF SEQ ID NOS: 343
     18 <170> SOFTWARE: PatentIn version 3.0
ERRORED SEQUENCES
     6041 <210> SEQ ID NO: 303
     6042 <211> LENGTH: (119) /09 slown
     6043 <212> TYPE: DNA
   > 6044 <213> ORGANISM: Artificial
     6046 <220> FEATURE:
     6047 <223> OTHER INFORMATION: synthetic construct
     6049 <220> FEATURE:
                                                - insert a hard return after 74007 303
   ♦ 6050 <221> NAME/KEY: SYNTHETIC GENE
     6051 <222> LOCATION: (1)..(119)
     6053 <400> SEQUENCE: 303 aggtcgtcac teegaegetg tttteacega caactacaeg egtetgegta (60
E--> 6055 aacagatggc tgcaaagaaa tacctgaact ccatcaagaa cgtgcgttaa tgagaattc
     6508 <210> SEQ ID NO: 329<211> 120
                              Tinsert a
    6509 <212> TYPE: DNA
⊕₩> 6510 <213> ORGANISM: Artificial
     6512 <220> FEATURE:
     6513 <223> OTHER INFORMATION: synthetic construct
     6515 <220> FEATURE:

✓> 6516 <221> NAME/KEY: SYNTHETIC GENE

     6517 <222> LOCATION: (1)..(120)
     6519 <400> SEQUENCE: 329
     6520 ggatecateg aaggtegtea eteegaeget gtttteaeeg acaactaeae gegtetgegt
                                                                                60
(€)4-> 6522 aaacagatgg ctgcaaagaa atacctgaac tccatcccga accgtatcta atgagaattc
                                                                               120
     6759 <210> SEQ ID NO: 343
     6760 <211> LENGTH: 120
                                     see P.Z
     6761 <212> TYPE: DNA
Q > 6762 <213 > ORGANISM: Artificial
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RAW SEQUENCE LISTING

DATE: 06/11/2002

PATENT APPLICATION: US/09/671,773

TIME: 14:12:21

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\06112002\1671773.raw

6764 <220> FEATURE:

6765 <223> OTHER INFORMATION: synthetic construct

6767 <220> FEATURE:

₩ > 6768 <221> NAME/KEY: SYNTHETIC GENE

6769 <222> LOCATION: (1)..(120)

6771 <400> SEQUENCE: 343

6772 ggatccatcg aaggtcgtca ctccgacgct gttttcaccg acaactacac gcgtctgcgt

6774 aaacagatgg cttccaagaa atacctgaac tccatccgta accgtatcta atgagaattc

E--> 6778/1

Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/11/2002 PATENT APPLICATION: US/09/671,773 TIME: 14:12:22

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\06112002\1671773.raw

## Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27 Seq#:28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51 Seq#:52,53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75 Seq#:76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99 Seq#:100,101,102,103,104,105,106,107,108,109,110,111,112,113,114,115,116,117 Seq#:118,119,120,121,122,123,124,125,126,127,128,129,130,131,132,133,134,135 Seq#:136,137,138,139,140,141,142,143,144,145,146,147,148,149,150,151,152,153 Seq#:154,155,156,157,158,159,160,161,162,163,164,165,166,167,168,169,170,171 Seq#:172,173,174,175,176,177,178,179,180,181,182,183,184,185,186,187,188,189 Seq#:190,191,192,193,194,195,196,197,198,199,200,201,202,203,204,205,206,207 Seq#:208,209,210,211,212,213,214,215,216,217,218,219,220,221,222,223,224,225 Seq#: 226,227,228,229,230,231,232,233,234,235,236,237,238,239,240,241,242,243 Seq#: 244,245,246,247,248,249,250,251,252,253,254,255,256,257,258,259,260,261 Seq#: 262, 263, 264, 265, 266, 267, 268, 269, 270, 271, 272, 273, 274, 275, 276, 277, 278, 279 $\mathtt{Seq\#:280,281,282,283,284,285,286,287,288,289,290,291,292,293,294,295,296,297}$ Seq#:298,299,300,301,302,303,304,305,306,307,308,309,310,311,312,313,314,315 Seq#:316,317,318,319,320,321,322,323,324,325,326,327,328,329,330,331,332,333 Seq#:334,335,336,337,338,339,340,341,342,343

 VERIFICATION SUMMARY
 DATE: 06/11/2002

 PATENT APPLICATION: US/09/671,773
 TIME: 14:12:22

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\06112002\1671773.raw

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:23 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:45 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:70 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:92 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:117 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:16
L:151 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:179 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:201 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:221 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:16
L:259 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:279 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:301 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:321 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:341 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:361 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:381 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:403 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:423 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:446 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:466 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20
L:486 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
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L:566 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:586 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
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L:626 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28
L:646 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:666 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30
L:686 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
L:706 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32
L:726 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33
L:746 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:34
L:766 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:35 L:786 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:36
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L:866 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:40
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L:906 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:42
L:926 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:43
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/671,773

DATE: 06/11/2002

TIME: 14:12:22

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\06112002\1671773.raw

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L:986\ M:220\ C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:46
L:1006 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:47
L:1026 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:48
L:1046 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:49
L:1066 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50
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L:1172\ M:257\ W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:55
L:1190~M:257~W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:56
L:3359 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:164
L:4576 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:222
L:4596 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:223
L:4614 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:224
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L:4692 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:228
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L:5204 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:256
L:5222 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:257
L:5240 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:258
L:5258 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:259
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## VERIFICATION SUMMARY

DATE: 06/11/2002 PATENT APPLICATION: US/09/671,773 TIME: 14:12:22

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\06112002\1671773.raw

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L:5348 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:264
L:5366 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:265
L:5384 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:266
L:5402 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:267
L:6055 M:254 E: No. of Bases conflict, LENGTH:Input:119 Counted:59 SEQ:303
L:6055 M:252 E: No. of Seq. differs, <211> LENGTH:Input:119 Found:59 SEQ:303
L:6519 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:6522 M:252 E: No. of Seq. differs, <211> LENGTH:Input:0 Found:120 SEQ:329
L:6778 M:254 E: No. of Bases conflict, this line has no nucleotides.
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